

Refine Search

Search Results -

Terms	Documents
L1 and (olson or Burvenich or Sydow or Zhang or Anvret).in.	24

Database:

US Pre-Grant Publication Full-Text Database
 US Patents Full-Text Database
 US OCR Full-Text Database
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Search:

L5

Search History

DATE: Monday, May 01, 2006 [Printable Copy](#) [Create Case](#)

<u>Set</u> <u>Name</u> side by side	<u>Query</u>	<u>Hit</u> <u>Count</u>	<u>Set</u> <u>Name</u> result set
<i>DB=PGPB,USPT,EPAB,JPAB,DWPI; PLUR=YES; OP=OR</i>			
<u>L5</u>	L1 and (olson or Burvenich or Sydow or Zhang or Anvret).in.	24	<u>L5</u>
<u>L4</u>	L3 and adh7\$4	7	<u>L4</u>
<u>L3</u>	L2 and parkinso\$4	120	<u>L3</u>
<u>L2</u>	L1 and (allel\$4 or polymorph\$6)	546	<u>L2</u>
<u>L1</u>	adh7\$6 or (alcoh\$4 same dehydrogenas\$4 same (huma\$4 or sapien\$4))	970	<u>L1</u>

END OF SEARCH HISTORY

Hit List

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Search Results - Record(s) 1 through 7 of 7 returned.

☐ 1. Document ID: US 20040241798 A1

Using default format because multiple data bases are involved.

L4: Entry 1 of 7

File: PGPB

Dec 2, 2004

PGPUB-DOCUMENT-NUMBER: 20040241798

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20040241798 A1

TITLE: ADH7 nucleotides

PUBLICATION-DATE: December 2, 2004

INVENTOR-INFORMATION:

NAME	CITY	STATE	COUNTRY
Olson, Lars	Lidingo		SE
Burvenich, Silvia	Danderyd		SE
Sydow, Olof	Bromma		SE
Anvret, Maria	Jarfalla		SE
Zhang, Zhiping	Huddinge		SE

US-CL-CURRENT: [435/69.1](#); [435/189](#), [435/320.1](#), [435/325](#), [435/6](#), [536/23.2](#)

Full	Title	Citation	Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KWIC	Draw D
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☐ 2. Document ID: US 20040214213 A1

L4: Entry 2 of 7

File: PGPB

Oct 28, 2004

PGPUB-DOCUMENT-NUMBER: 20040214213

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20040214213 A1

TITLE: ADH1C

PUBLICATION-DATE: October 28, 2004

INVENTOR-INFORMATION:

NAME	CITY	STATE	COUNTRY
Burvenich, Silvia	Washington	DC	US
Carmine, Andrea	Bromma		SE
Galter, Dagmar	Bromma		SE

US-CL-CURRENT: 435/6

Full	Title	Citation	Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KMC	Drawn	Drawn Date
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Sep 2, 2004

US-CL-CURRENT: 435/6; 530/350, 536/24.3

Full	Title	Citation	Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	QWIC	Draw D
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Jun 10, 2004

US-CL-CURRENT: 514/44

Full	Title	Citation	Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KMIC	Draw D
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☐ 5. Document ID: US 20010034023 A1

L4: Entry 5 of 7

File: PGPB

Oct 25, 2001

PGPUB-DOCUMENT-NUMBER: 20010034023

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20010034023 A1

TITLE: Gene sequence variations with utility in determining the treatment of disease, in genes relating to drug processing

PUBLICATION-DATE: October 25, 2001

INVENTOR-INFORMATION:

NAME	CITY	STATE	COUNTRY
Stanton, Vincent P. JR.	Belmont	MA	US
Zillmann, Martin	Shrewsbury	MA	US

US-CL-CURRENT: 435/6; 702/20

Full	Title	Citation	Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KWIC	Draw D
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☐ 6. Document ID: US 6673549 B1

L4: Entry 6 of 7

File: USPT

Jan 6, 2004

US-PAT-NO: 6673549

DOCUMENT-IDENTIFIER: US 6673549 B1

TITLE: Genes expressed in C3A liver cell cultures treated with steroids

DATE-ISSUED: January 6, 2004

INVENTOR-INFORMATION:

NAME	CITY	STATE	ZIP CODE	COUNTRY
Furness; L. Michael	Suffolk			GB
Buchbinder; Jenny L.	San Francisco	CA		

US-CL-CURRENT: 435/6; 435/287.2, 435/7.1, 514/44, 536/23.1

Full	Title	Citation	Front	Review	Classification	Date	Reference			Claims	KWIC	Draw D
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☐ 7. Document ID: WO 200224958 A1, AU 200196276 A

L4: Entry 7 of 7

File: DWPI

Mar 28, 2002

DERWENT-ACC-NO: 2002-352009

DERWENT-WEEK: 200252

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TITLE: New genetic variants of the alcohol dehydrogenase 7 (class IV) mu or sigma polypeptide (ADH7) gene having polymorphisms, useful for treating disorders

affected by ADH7 isogene expression or function, e.g. cancer or Parkinson's disease

INVENTOR: BIEGLECKI, K M; FINKEL, K ; KAZEMI, A ; KOSHY, B ; PARKS, K E ; SAUSKER, E A

PRIORITY-DATA: 2000US-233520P (September 19, 2000)

PATENT-FAMILY:

PUB-NO	PUB-DATE	LANGUAGE	PAGES	MAIN-IPC
<u>WO 200224958 A1</u>	March 28, 2002	E	114	C12Q001/68
<u>AU 200196276 A</u>	April 2, 2002		000	C12Q001/68

INT-CL (IPC): C07 H 21/04; C12 Q 1/68

Full	Title	Citation	Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KWIC	Draw D
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Terms	Documents
L3 and adh7\$4	7

Display Format:

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=> d his full

(FILE 'HOME' ENTERED AT 11:20:59 ON 01 MAY 2006)

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 11:21:25 ON 01 MAY 2006
SEA ADH?? OR (ALCOHOL?(S)DEHYDROGENAS?(S)(HUMA? OR SAPIEN?))

7 FILE ADISCTI
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886 FILE USPATFULL
69 FILE USPAT2
1 FILE WATER
91 FILE WPIDS
1 FILE WPIFV
91 FILE WPINDEX
8 FILE IPA
1 FILE NAPRALERT
3 FILE NLDB

L1 QUE ADH?? OR (ALCOHOL?(S) DEHYDROGENAS?(S)(HUMA? OR SAPIEN?))

D RANK

FILE 'BIOSIS, TOXCENTER, USPATFULL, CAPLUS, MEDLINE, SCISEARCH, EMBASE,
BIOTECHNO, ESBIODBASE, PASCAL, LIFESCI' ENTERED AT 11:25:52 ON 01 MAY 2006

L2 7283 SEA ADH?? OR (ALCOHOL?(S) DEHYDROGENAS?(S)(HUMA? OR SAPIEN?))

L3 1582 SEA L2 AND (ALLELE? OR POLYMORPH?)

L4 127 SEA L3 AND PARKINSON?

L5 117 DUP REM L4 (10 DUPLICATES REMOVED)

D TI L5 1-117

D IBIB ABS L5 107 105 99 80 27 25 9

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NEWS	1		Web Page URLs for STN Seminar Schedule - N. America
NEWS	2		"Ask CAS" for self-help around the clock
NEWS	3	DEC 23	New IPC8 SEARCH, DISPLAY, and SELECT fields in USPATFULL/ USPAT2
NEWS	4	JAN 13	IPC 8 searching in IFIPAT, IFIUDb, and IFICDB
NEWS	5	JAN 13	New IPC 8 SEARCH, DISPLAY, and SELECT enhancements added to INPADOC
NEWS	6	JAN 17	Pre-1988 INPI data added to MARPAT
NEWS	7	JAN 17	IPC 8 in the WPI family of databases including WPIFV
NEWS	8	JAN 30	Saved answer limit increased
NEWS	9	FEB 21	STN AnaVist, Version 1.1, lets you share your STN AnaVist visualization results
NEWS	10	FEB 22	The IPC thesaurus added to additional patent databases on STN
NEWS	11	FEB 22	Updates in EPFULL; IPC 8 enhancements added
NEWS	12	FEB 27	New STN AnaVist pricing effective March 1, 2006
NEWS	13	FEB 28	MEDLINE/LMEDLINE reload improves functionality
NEWS	14	FEB 28	TOXCENTER reloaded with enhancements
NEWS	15	FEB 28	REGISTRY/ZREGISTRY enhanced with more experimental spectral property data
NEWS	16	MAR 01	INSPEC reloaded and enhanced
NEWS	17	MAR 03	Updates in PATDPA; addition of IPC 8 data without attributes
NEWS	18	MAR 08	X.25 communication option no longer available after June 2006
NEWS	19	MAR 22	EMBASE is now updated on a daily basis
NEWS	20	APR 03	New IPC 8 fields and IPC thesaurus added to PATDPAFULL
NEWS	21	APR 03	Bibliographic data updates resume; new IPC 8 fields and IPC thesaurus added in PCTFULL
NEWS	22	APR 04	STN AnaVist \$500 visualization usage credit offered
NEWS	23	APR 12	LINSPEC, learning database for INSPEC, reloaded and enhanced
NEWS	24	APR 12	Improved structure highlighting in FQHIT and QHIT display in MARPAT
NEWS	25	APR 12	Derwent World Patents Index to be reloaded and enhanced during second quarter; strategies may be affected
NEWS EXPRESS			FEBRUARY 15 CURRENT VERSION FOR WINDOWS IS V8.01a, CURRENT MACINTOSH VERSION IS V6.0c(ENG) AND V6.0Jc(JP), AND CURRENT DISCOVER FILE IS DATED 19 DECEMBER 2005. V8.0 AND V8.01 USERS CAN OBTAIN THE UPGRADE TO V8.01a AT http://download.cas.org/express/v8.0-Discover/
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FILE 'HOME' ENTERED AT 11:20:59 ON 01 MAY 2006

=> index bioscience medicine

FILE 'DRUGMONOG' ACCESS NOT AUTHORIZED

COST IN U.S. DOLLARS

SINCE FILE	TOTAL
ENTRY	SESSION
0.21	0.21

FULL ESTIMATED COST

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 11:21:25 ON 01 MAY 2006

71 FILES IN THE FILE LIST IN STNINDEX

Enter SET DETAIL ON to see search term postings or to view search error messages that display as 0* with SET DETAIL OFF.

=> s adh?? or (alcohol?(s)dehydrogenas?(s)(huma? or sapien?))

7	FILE ADISCTI
5	FILE ADISINSIGHT
3	FILE ADISNEWS
33	FILE AGRICOLA
33	FILE ANABSTR
3	FILE ANTE
1	FILE AQUALINE
9	FILE AQUASCI
34	FILE BIOENG
1281	FILE BIOSIS
165	FILE BIOTECHABS
165	FILE BIOTECHDS
463	FILE BIOTECHNO

13 FILES SEARCHED...

86	FILE CABA
761	FILE CAPLUS
19	FILE CEABA-VTB
2	FILE CIN
31	FILE CONFSCI
2	FILE CROPU
55	FILE DDFB
57	FILE DDFU

2317 FILE DGENE
 23 FILES SEARCHED...
 52 FILE DISSABS
 55 FILE DRUGB
 78 FILE DRUGU
 6 FILE EMBAL
 518 FILE EMBASE
 440 FILE ESBIODBASE
 10 FILE FROSTI
 15 FILE FSTA
 1688 FILE GENBANK
 35 FILES SEARCHED...
 7 FILE HEALSAFE
 44 FILE IFIPAT
 54 FILE JICST-EPLUS
 6 FILE KOSMET
 364 FILE LIFESCI
 600 FILE MEDLINE
 7 FILE NTIS
 2 FILE OCEAN
 424 FILE PASCAL
 48 FILES SEARCHED...
 1 FILE PHIN
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 2 FILE PROUSDDR
 3 FILE RDISCLOSURE
 591 FILE SCISEARCH
 955 FILE TOXCENTER
 60 FILES SEARCHED...
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 69 FILE USPAT2
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 91 FILE WPIDS
 1 FILE WPIFV
 91 FILE WPINDEX
 8 FILE IPA
 1 FILE NAPRALERT
 3 FILE NLDB

55 FILES HAVE ONE OR MORE ANSWERS, 71 FILES SEARCHED IN STNINDEX

L1 QUE ADH7? OR (ALCOHOL?(S) DEHYDROGENAS?(S) (HUMA? OR SAPIEN?))

=> d rank

F1	2317	DGENE
F2	1688	GENBANK
F3	1281	BIOSIS
F4	955	TOXCENTER
F5	886	USPATFULL
F6	761	CAPLUS
F7	600	MEDLINE
F8	591	SCISEARCH
F9	518	EMBASE
F10	463	BIOTECHNO
F11	440	ESBIODBASE
F12	424	PASCAL
F13	364	LIFESCI
F14	165	BIOTECHABS
F15	165	BIOTECHDS
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F20	69	USPAT2
F21	57	DDFU

F22	55	DDFB
F23	55	DRUGB
F24	54	JICST-EPLUS
F25	52	DISSABS
F26	44	IFIPAT
F27	34	BIOENG
F28	33	AGRICOLA
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F30	31	CONFSCI
F31	19	CEABA-VTB
F32	19	PROMT
F33	15	FSTA
F34	10	FROSTI
F35	9	AQUASCI
F36	8	IPA
F37	7	ADISCTI
F38	7	HEALSAFE
F39	7	NTIS
F40	6	EMBAL
F41	6	KOSMET
F42	5	ADISINSIGHT
F43	3	ADISNEWS
F44	3	ANTE
F45	3	RDISCLOSURE
F46	3	NLDB
F47	2	CIN
F48	2	CROPU
F49	2	OCEAN
F50	2	PROUSDDR
F51	1	AQUALINE
F52	1	PHIN
F53	1	WATER
F54	1	WPIFV
F55	1	NAPRALERT

=> file f3-f13

COST IN U.S. DOLLARS

SINCE FILE

ENTRY

TOTAL

SESSION

FULL ESTIMATED COST

4.27

4.48

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=> s adh7? or (alcohol?(s)dehydrogenas?(s)(huma? or sapien?))

9 FILES SEARCHED...

L2 7283 ADH7? OR (ALCOHOL?(S) DEHYDROGENAS?(S) (HUMA? OR SAPIEN?))

=> s l2 and (allele? or polymorph?)

L3 1582 L2 AND (ALLELE? OR POLYMORPH?)

=> s l3 and parkinson?

L4 127 L3 AND PARKINSON?

=> dup rem l4

PROCESSING COMPLETED FOR L4

L5 117 DUP REM L4 (10 DUPLICATES REMOVED)

=> d ti l5 1-117

L5 ANSWER 1 OF 117 USPATFULL on STN

TI Novel 13237, 18480, 2245, 16228, 7677, 26320, 46619, 33166, 16836,
46867, 21617, 55562, 39228, 62088, 46745, 23155, 21657, 42755, 32229,
22325, 46863 and 32252 molecules and uses therefor

L5 ANSWER 2 OF 117 USPATFULL on STN

TI Proteins, polynucleotides encoding them and methods of using the same

L5 ANSWER 3 OF 117 USPATFULL on STN

TI Drug targets for the treatment of neurodegenerative disorders

L5 ANSWER 4 OF 117 USPATFULL on STN

TI Drug targets for the treatment of neurodegenerative disorders

L5 ANSWER 5 OF 117 USPATFULL on STN

TI Diagnosis and treatment system for "reward deficiency syndrome" (RDS)
and related behaviors

L5 ANSWER 6 OF 117 USPATFULL on STN

TI Compositions and treatment methods

L5 ANSWER 7 OF 117 USPATFULL on STN

TI Methods and compositions for treating hematological disorders using
9118, 990, 17662, 81982, 630, 21472, 17692, 19290, 21620, 21689, 28899,
53659, 64549, 9465, 23544, 7366, 27417, 57259, 21844, 943, 2061, 5891,
9137, 13908, 14310, 17600, 25584, 27824, 28469, 38947, 53003, 965,
56639, 9661, 16052, 1521, 6662, 13913, 12405 or 5014

L5 ANSWER 8 OF 117 USPATFULL on STN

TI Molecular toxicology modeling

L5 ANSWER 9 OF 117 USPATFULL on STN

TI Cloning of a gene mutation for parkinson's disease

L5 ANSWER 10 OF 117 USPATFULL on STN

TI Intracellular amyloid-beta binding (ERAB) polypeptide-related assays
 L5 ANSWER 11 OF 117 USPATFULL on STN
 TI Substituted furo[2,3-b]pyridine derivatives
 L5 ANSWER 12 OF 117 USPATFULL on STN
 TI Flea head, nerve cord, hindgut and malpighian tubule nucleic acid molecules, proteins and uses thereof
 L5 ANSWER 13 OF 117 USPATFULL on STN
 TI Novel nucleic acids and polypeptides
 L5 ANSWER 14 OF 117 USPATFULL on STN
 TI Novel human genes and methods of use thereof
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 TI Enzymes
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 TI Novel method for down-regulation of amyloid
 L5 ANSWER 17 OF 117 USPATFULL on STN
 TI Therapeutic treatment methods 2
 L5 ANSWER 18 OF 117 USPATFULL on STN
 TI Pharmaceutical compositions and methods for treating, preventing, and managing cholesterol, dyslipidemia, and related disorders
 L5 ANSWER 19 OF 117 USPATFULL on STN
 TI Method of treating cancer using dithiocarbamate derivatives
 L5 ANSWER 20 OF 117 USPATFULL on STN
 TI 87 human secreted proteins
 L5 ANSWER 21 OF 117 USPATFULL on STN
 TI Neurothrophic componentss of the adnf 1 complex
 L5 ANSWER 22 OF 117 USPATFULL on STN
 TI Methods and compositions for treating hematological disorders using 9118, 990, 17662, 81982, 630, 21472, 17692, 19290, 21620, 21689, 28899, 53659, 64549, 9465, 23544, 7366, 27417, 57259, 21844, 943, 2061, 5891, 9137, 13908, 14310, 17600, 25584, 27824, 28469, 38947, 53003, 965, 56639, 9661, 16052, 1521, 6662, 13913, 12405 or 5014
 L5 ANSWER 23 OF 117 USPATFULL on STN
 TI Compositions and methods for modulating S-nitrosoglutathione reductase
 L5 ANSWER 24 OF 117 USPATFULL on STN
 TI Diagnosis and treatment system for reward deficiency syndrome (RDS) and related behaviors
 L5 ANSWER 25 OF 117 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN
 TI A rare truncating mutation in ADH1C (G78Stop) shows significant association with Parkinson disease in a large international sample. DUPLICATE 1
 L5 ANSWER 26 OF 117 USPATFULL on STN
 TI ADH1C DUPLICATE 2
 L5 ANSWER 27 OF 117 USPATFULL on STN
 TI ADH7 nucleotides
 L5 ANSWER 28 OF 117 USPATFULL on STN
 TI Method for cohort selection

L5 ANSWER 29 OF 117 USPATFULL on STN
TI Gene sequence variations with utility in determining the treatment of disease, in genes relating to drug processing

L5 ANSWER 30 OF 117 USPATFULL on STN
TI Therapeutic treatment methods

L5 ANSWER 31 OF 117 USPATFULL on STN
TI Novel human enzyme family members and uses thereof

L5 ANSWER 32 OF 117 USPATFULL on STN
TI Method of treating neurological diseases

L5 ANSWER 33 OF 117 USPATFULL on STN
TI Drug metabolizing enzymes

L5 ANSWER 34 OF 117 USPATFULL on STN
TI Molecular toxicology modeling

L5 ANSWER 35 OF 117 USPATFULL on STN
TI Novel proteins and nucleic acids encoding same

L5 ANSWER 36 OF 117 USPATFULL on STN
TI Novel proteins and nucleic acids encoding same

L5 ANSWER 37 OF 117 USPATFULL on STN
TI Novel proteins and nucleic acids encoding same

L5 ANSWER 38 OF 117 USPATFULL on STN
TI Novel 13237, 18480, 2245, 16228, 7677, 26320, 46619, 33166, 16836, 46867, 21617, 55562, 39228, 62088, 46745, 23155, 21657, 42755, 32229, 22325, 46863 and 32252 molecules and uses therefor

L5 ANSWER 39 OF 117 USPATFULL on STN
TI Furanone derivatives

L5 ANSWER 40 OF 117 USPATFULL on STN
TI Novel proteins and nucleic acids encoding same

L5 ANSWER 41 OF 117 USPATFULL on STN
TI Proteins and nucleic acids encoding same

L5 ANSWER 42 OF 117 USPATFULL on STN
TI Novel proteins and nucleic acids encoding same

L5 ANSWER 43 OF 117 USPATFULL on STN
TI Novel proteins and nucleic acids encoding same

L5 ANSWER 44 OF 117 USPATFULL on STN
TI Novel human polynucleotides and polypeptides encoded thereby

L5 ANSWER 45 OF 117 USPATFULL on STN
TI Novel 26199, 33530, 33949, 47148, 50226, 58764, 62113, 32144, 32235, 23565, 13305, 14911, 86216, 25206 and 8843 molecules and uses therefor

L5 ANSWER 46 OF 117 USPATFULL on STN
TI Proteins, polynucleotides encoding them and methods of using the same

L5 ANSWER 47 OF 117 USPATFULL on STN
TI Nucleic acids and polypeptides

L5 ANSWER 48 OF 117 USPATFULL on STN
TI Genes expressed in C3A liver cell cultures treated with steroids

L5 ANSWER 49 OF 117 USPATFULL on STN
 TI Nucleic acids, proteins, and antibodies

L5 ANSWER 50 OF 117 USPATFULL on STN
 TI Methods for genetic analysis of DNA using biased amplification of **polymorphic** sites

L5 ANSWER 51 OF 117 USPATFULL on STN
 TI Novel nucleic acids and polypeptides

L5 ANSWER 52 OF 117 USPATFULL on STN
 TI Novel nucleic acids and polypeptides

L5 ANSWER 53 OF 117 USPATFULL on STN
 TI 207 human secreted proteins

L5 ANSWER 54 OF 117 USPATFULL on STN
 TI Furanone derivatives

L5 ANSWER 55 OF 117 USPATFULL on STN
 TI Individualization of therapy with antipsychotics

L5 ANSWER 56 OF 117 USPATFULL on STN
 TI Nucleic acid hairpin probes and uses thereof

L5 ANSWER 57 OF 117 USPATFULL on STN
 TI Novel method for down-regulation of amyloid

L5 ANSWER 58 OF 117 USPATFULL on STN
 TI Novel human genes and methods of use thereof

L5 ANSWER 59 OF 117 USPATFULL on STN
 TI Sequence-directed DNA-binding molecules compositions and methods

L5 ANSWER 60 OF 117 USPATFULL on STN
 TI Human oxidoreductase proteins

L5 ANSWER 61 OF 117 USPATFULL on STN
 TI Novel nucleic acids and polypeptides

L5 ANSWER 62 OF 117 USPATFULL on STN
 TI Novel human enzyme family members and uses thereof

L5 ANSWER 63 OF 117 USPATFULL on STN
 TI Blood cell deficiency treatment method

L5 ANSWER 64 OF 117 USPATFULL on STN
 TI Methods for genetic analysis of DNA to detect sequence variances

L5 ANSWER 65 OF 117 USPATFULL on STN
 TI Flea head, nerve cord, hindgut and malpighian tubule nucleic acid molecules, proteins and uses thereof

L5 ANSWER 66 OF 117 USPATFULL on STN
 TI Restriction enzyme genotyping

L5 ANSWER 67 OF 117 USPATFULL on STN
 TI Individualization of therapy with antidepressants

L5 ANSWER 68 OF 117 USPATFULL on STN
 TI 87 human secreted proteins

L5 ANSWER 69 OF 117 USPATFULL on STN
 TI Nucleic acid molecules, polypeptides and uses therefor, including diagnosis and treatment of Alzheimer's disease

L5 ANSWER 70 OF 117 USPATFULL on STN
TI Secreted protein HFEAF41

L5 ANSWER 71 OF 117 USPATFULL on STN
TI Novel human 39228, 21956, 25856, 22244, 8701, 32263, 50250, 55158, 47765, 62088, 50566, and 48118 molecules and uses therefor

L5 ANSWER 72 OF 117 USPATFULL on STN
TI Nucleic acids, proteins, and antibodies

L5 ANSWER 73 OF 117 USPATFULL on STN
TI Secreted protein HFEAF41

L5 ANSWER 74 OF 117 USPATFULL on STN
TI Novel nucleic acid sequences encoding a human ubiquitin protease, lipase, dynamin, short chain dehydrogenase, and ADAM-TS metalloprotease and uses therefor

L5 ANSWER 75 OF 117 USPATFULL on STN
TI Computer-assisted means for assessing lifestyle risk factors

L5 ANSWER 76 OF 117 USPATFULL on STN
TI 33200, a novel human dehydrogenase/reductase family member and uses thereof

L5 ANSWER 77 OF 117 USPATFULL on STN
TI Secreted protein HFEAF41

L5 ANSWER 78 OF 117 USPATFULL on STN
TI Nucleic acids and polypeptides

L5 ANSWER 79 OF 117 USPATFULL on STN
TI 32140, a novel human aldehyde dehydrogenase and uses therefor

L5 ANSWER 80 OF 117 SCISEARCH COPYRIGHT (c) 2006 The Thomson Corporation on STN
TI Frequencies of single nucleotide **polymorphism** in alcohol dehydrogenase7 gene in patients with multiple system atrophy and controls

L5 ANSWER 81 OF 117 USPATFULL on STN
TI Human polynucleotides, polypeptides, and antibodies

L5 ANSWER 82 OF 117 USPATFULL on STN
TI Novel method for down-regulation of amyloid

L5 ANSWER 83 OF 117 USPATFULL on STN
TI Secreted protein HFEAF41

L5 ANSWER 84 OF 117 USPATFULL on STN
TI Nucleic acids, proteins, and antibodies

L5 ANSWER 85 OF 117 USPATFULL on STN
TI 25206, a novel human short-chain dehydrogenase/reductase family member and uses thereof

L5 ANSWER 86 OF 117 USPATFULL on STN
TI Human polynucleotides, polypeptides, and antibodies

L5 ANSWER 87 OF 117 USPATFULL on STN
TI Nucleic acid hairpin probes and uses thereof

L5 ANSWER 88 OF 117 USPATFULL on STN
TI 25219, a novel human aminotransferase and uses therefor

L5 ANSWER 89 OF 117 USPATFULL on STN
 TI Nucleic acids, proteins, and antibodies

L5 ANSWER 90 OF 117 USPATFULL on STN
 TI 22244 and 8701, novel human dehydrogenases and uses thereof

L5 ANSWER 91 OF 117 USPATFULL on STN
 TI Nucleic acids, proteins and antibodies

L5 ANSWER 92 OF 117 USPATFULL on STN
 TI Nucleic acids, proteins and antibodies

L5 ANSWER 93 OF 117 USPATFULL on STN
 TI Nucleic, acids, proteins, and antibodies

L5 ANSWER 94 OF 117 USPATFULL on STN
 TI 21612, 21615, 21620, 21676, 33756, novel human alcohol dehydrogenases

L5 ANSWER 95 OF 117 USPATFULL on STN
 TI 21509 and 33770, novel human dehydrogenase family members and uses thereof

L5 ANSWER 96 OF 117 USPATFULL on STN
 TI Methods for genetic analysis of DNA using biased amplification of polymorphic sites

L5 ANSWER 97 OF 117 USPATFULL on STN
 TI Sequence directed DNA binding molecules compositions and methods

L5 ANSWER 98 OF 117 BIOTECHNO COPYRIGHT 2006 Elsevier Science B.V. on STN
 TI Common psychiatric diseases and human genetic variation

L5 ANSWER 99 OF 117 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN
 TI ADH1C NONSENSE MUTATION IN PARKINSON DISEASE. DUPLICATE 3

L5 ANSWER 100 OF 117 USPATFULL on STN
 TI Gene sequence variations with utility in determining the treatment of disease, in genes relating to drug processing

L5 ANSWER 101 OF 117 USPATFULL on STN
 TI Intracellular amyloid-beta peptide binding (ERAB) polypeptide

L5 ANSWER 102 OF 117 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN
 TI Alcohol dehydrogenase polymorphism and Parkinson's disease. DUPLICATE 4

L5 ANSWER 103 OF 117 USPATFULL on STN
 TI Surrogate tolerogenesis for the development of tolerance to xenografts

L5 ANSWER 104 OF 117 USPATFULL on STN
 TI Sequence-directed DNA binding molecules compositions and methods

L5 ANSWER 105 OF 117 MEDLINE on STN
 TI Alcohol dehydrogenase alleles in Parkinson's disease. DUPLICATE 5

L5 ANSWER 106 OF 117 CAPLUS COPYRIGHT 2006 ACS on STN
 TI The emerging field of ecogenetics

L5 ANSWER 107 OF 117 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN
 TI Association of an alcohol dehydrogenase allele with Parkinson's disease.

L5 ANSWER 108 OF 117 USPATFULL on STN
 TI Method of determining DNA sequence preference of a DNA-binding molecule

L5 ANSWER 109 OF 117 USPATFULL on STN
 TI Sequence-directed DNA-binding molecules compositions and methods

L5 ANSWER 110 OF 117 USPATFULL on STN
 TI Sequence-directed DNA-binding molecules compositions and methods

L5 ANSWER 111 OF 117 USPATFULL on STN
 TI Screening assay for the detection of DNA-binding molecules

L5 ANSWER 112 OF 117 USPATFULL on STN
 TI Method of constructing sequence-specific DNA-binding molecules

L5 ANSWER 113 OF 117 USPATFULL on STN
 TI Method of ordering sequence binding preferences of a DNA-binding molecule

L5 ANSWER 114 OF 117 USPATFULL on STN
 TI Sequence-directed DNA-binding molecules compositions and methods

L5 ANSWER 115 OF 117 USPATFULL on STN
 TI Allelic diagnosis of susceptibility to compulsive disorder

L5 ANSWER 116 OF 117 USPATFULL on STN
 TI Allelic association of the human dopamine(D2) receptor gene in compulsive disorders

L5 ANSWER 117 OF 117 USPATFULL on STN
 TI Allelic association of the human dopamine (D.sub.2) receptor gene in compulsive disorders such as alcoholism

=> d ibib abs 15 107 105 99 80 27 25 9

L5 ANSWER 107 OF 117 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN

ACCESSION NUMBER: 2001:97716 BIOSIS
 DOCUMENT NUMBER: PREV200100097716
 TITLE: Association of an alcohol dehydrogenase allele with **Parkinson's** disease.
 AUTHOR(S): Buervenich, S. [Reprint author]; Sydow, O.; Carmine, A.; Galter, D.; Zhang, Z.; Anvret, M.; Olson, L.
 CORPORATE SOURCE: Danderyd Hospital, Stockholm, Sweden
 SOURCE: Society for Neuroscience Abstracts, (2000) Vol. 26, No. 1-2, pp. Abstract No.-476.10. print.
 Meeting Info.: 30th Annual Meeting of the Society of Neuroscience. New Orleans, LA, USA. November 04-09, 2000. Society for Neuroscience.
 ISSN: 0190-5295.
 DOCUMENT TYPE: Conference; (Meeting)
 Conference; Abstract; (Meeting Abstract)
 LANGUAGE: English
 ENTRY DATE: Entered STN: 21 Feb 2001
 Last Updated on STN: 15 Feb 2002

AB Because of the important role of **alcohol dehydrogenases** (ADH; EC 1.1.1.1) in retinoid and dopamine metabolism and/or aldehyde detoxification, mutations in genes for these enzymes, clustered on **human** chromosome 4q21-25, may be genetic risk factors for **Parkinson's** disease and other disorders of the dopamine system. Recently, a form of autosomal-dominant **parkinsonism** has been mapped to this area and a mutation was found in the gene for synuclein, which is mapped in close vicinity to the alcohol dehydrogenase gene

cluster. Because of the lack of synuclein mutations in other large **Parkinson's** disease materials, the possibility of further genetic factors in the linked region on chromosome four remains. We sequenced the human class IV ADH gene (previously named **ADH7**, currently termed **ADH4** according to the most recent class-oriented nomenclature) in ten **Parkinson** patients. Seven **polymorphisms** were identified. These **polymorphisms** could be assigned to four **alleles** (A1-A4) in our Swedish sample. Frequencies of those four **alleles** and the wildtype **allele** in 78 PD patients and 130 controls were determined. A significant association of the A1 **allele** with PD (odds ratio 2.87 (95% CI 1.35 to 6.08)) was found. The association was strongest in familial cases (odds ratio 4.86 (95% CI 1.89 to 12.75)). Thus, our results show an association between a certain **ADH4 allele** and PD. This suggests a role for genetic variations of **ADH4** as risk factors for the development of PD. Functional studies are currently carried out to elucidate the effects of the identified **polymorphisms** on **ADH4** gene function.

L5 ANSWER 105 OF 117 MEDLINE on STN DUPLICATE 5
 ACCESSION NUMBER: 2000452803 MEDLINE
 DOCUMENT NUMBER: PubMed ID: 11009184
 TITLE: Alcohol dehydrogenase **alleles** in **Parkinson's** disease.
 AUTHOR: Buervenich S; Sydow O; Carmine A; Zhang Z; Anvret M; Olson L
 CORPORATE SOURCE: Department of Neuroscience, Karolinska Institute, Stockholm, Sweden.
 SOURCE: Movement disorders : official journal of the Movement Disorder Society, (2000 Sep) Vol. 15, No. 5, pp. 813-8. Journal code: 8610688. ISSN: 0885-3185.
 PUB. COUNTRY: United States
 DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)
 LANGUAGE: English
 FILE SEGMENT: Priority Journals
 ENTRY MONTH: 200101
 ENTRY DATE: Entered STN: 22 Mar 2001
 Last Updated on STN: 22 Mar 2001
 Entered Medline: 9 Jan 2001

AB Mutations in alcohol dehydrogenase (ADH; EC 1.1.1.1) genes may be of interest in the etiology of **Parkinson's** disease (PD) because of the important role these enzymes play in retinoid and dopamine metabolism and/or aldehyde detoxification. The location of several alcohol dehydrogenase genes in a cluster on chromosome 4 lends further support to ADH genes being candidates for this disorder, because recently a form of autosomal-dominant **parkinsonism** has been mapped to this area. We sequenced the promoter and coding regions and part of the introns of the human class IV ADH gene in 10 patients with PD. Seven different **polymorphisms** were identified. These **polymorphisms** could be assigned to four **alleles** (A1-A4). We then determined the frequencies of those four **alleles** and the wild-type **allele** in 78 patients with PD and 130 control subjects and found a significant association of the A1 **allele** with PD (odds ratio = 2.87; 95% confidence interval = 1.35-6.08). In familial cases, the association was strongest (odds ratio = 4.86; 95% confidence interval = 1.89-12.75). Two patients were homozygous for A1 whereas none of the 130 control subjects was found to be homozygous. Our results show an association between a certain **ADH4** (formerly known as **ADH7** in humans) **allele** and PD. This suggests a role for genetic variations of **ADH4** as risk factors for the development of PD. Our data also show that the observed **polymorphisms** alone are not sufficient to cause symptoms. Further genetic and/or environmental factors have to be involved.

L5 ANSWER 99 OF 117 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN DUPLICATE 3

ACCESSION NUMBER: 2003:380970 BIOSIS
DOCUMENT NUMBER: PREV200300380970
TITLE: ADH1C NONSENSE MUTATION IN PARKINSON DISEASE.
AUTHOR(S): Buervenich, S. [Reprint Author]; Carmine, A. [Reprint Author]; Galter, D. [Reprint Author]; Matsuura, T.; Ashizawa, T.; Wullner, U.; Klockgether, T.; Nissbrandt, H.; Meister, B. [Reprint Author]; Anvret, M.; Sydow, O.; Olson, L. [Reprint Author]
CORPORATE SOURCE: Neuroscience, Molecular Medicine, Clinical Neuroscience, Karolinska Institutet, Stockholm, Sweden
SOURCE: Society for Neuroscience Abstract Viewer and Itinerary Planner, (2002) Vol. 2002, pp. Abstract No. 887.5. <http://sfn.scholarone.com.cd-rom>. Meeting Info.: 32nd Annual Meeting of the Society for Neuroscience. Orlando, Florida, USA. November 02-07, 2002. Society for Neuroscience.
DOCUMENT TYPE: Conference; (Meeting)
Conference; (Meeting Poster)
Conference; Abstract; (Meeting Abstract)
LANGUAGE: English
ENTRY DATE: Entered STN: 20 Aug 2003
Last Updated on STN: 20 Aug 2003

AB Genes encoding alcohol and aldehyde dehydrogenases may be considered candidate genes involved in the pathogenesis of neurodegenerative disorders due to the multiple roles these enzymes play in detoxification pathways, dopamine metabolism and retinoic acid synthesis. In a previous study, association of a certain allele of human class IV alcohol dehydrogenase (ADH4) with Parkinsons disease was found in our Swedish sample. Since the gene for ADH4 is located within a large cluster of alcohol dehydrogenases on human chromosome 4q21-q23, we extended the molecular genetic analyses to adjacent genes encoding other alcohol dehydrogenases. Thirteen known single-nucleotide polymorphisms (SNPs) distributed across this cluster were selected for further analysis in a subsequent study. The SNPs were located within the genes coding for ADH1B and ADH1C, class III ADH and class IV ADH. Distributions of these SNPs between 123 Swedish Parkinson patients and 127 geographically matched controls were investigated. One rare nonsense mutation in the gene encoding ADH1C was present in three patients with Parkinsons disease but completely absent from control samples. This mutation leads to an early truncation of the protein chain (after 78 of 375 amino acid residues). Neither the active site nor the cofactor binding site are included in the remaining protein chain.

L5 ANSWER 80 OF 117 SCISEARCH COPYRIGHT (c) 2006 The Thomson Corporation on STN

ACCESSION NUMBER: 2003:837973 SCISEARCH
THE GENUINE ARTICLE: 723PF
TITLE: Frequencies of single nucleotide polymorphism in alcohol dehydrogenase7 gene in patients with multiple system atrophy and controls
AUTHOR: Kim H S; Lee M S (Reprint)
CORPORATE SOURCE: Yonsei Univ, Yongdong Severance Hosp, Coll Med, Dept Neurol, Brain Korea Project 21, 146-92 Dogok Dong, Seoul 120749, South Korea (Reprint); Yonsei Univ, Yongdong Severance Hosp, Coll Med, Dept Neurol, Brain Korea Project 21, Seoul 120749, South Korea
COUNTRY OF AUTHOR: South Korea
SOURCE: MOVEMENT DISORDERS, (SEP 2003) Vol. 18, No. 9, pp. 1065-1067.
ISSN: 0885-3185.
PUBLISHER: WILEY-LISS, DIV JOHN WILEY & SONS INC, 605 THIRD AVE, NEW YORK, NY 10158-0012 USA.
DOCUMENT TYPE: Article; Journal

LANGUAGE: English

REFERENCE COUNT: 21

ENTRY DATE: Entered STN: 10 Oct 2003

Last Updated on STN: 10 Oct 2003

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS

AB A polymerase chain reaction and direct sequencing of the ADH7 gene were carried out in 50 controls and 50 patients with probable multiple system atrophy (MSA). Seven SNPs, one insertion, and one mismatch were found in patients with MSA and controls. There was no significant difference in the frequencies of each SNP between the patients and the controls ($P > 0.05$). Interpretation of this negative finding should be cautious in view of the relatively small number of cases. (C) 2003 Movement Disorder Society.

L5 ANSWER 27 OF 117 USPATFULL on STN

ACCESSION NUMBER: 2004:307127 USPATFULL

TITLE: ADH7 nucleotides

INVENTOR(S): Olson, Lars, Lidings, SWEDEN
Burvenich, Silvia, Danderyd, SWEDEN
Sydow, Olof, Bromma, SWEDEN
Anvret, Maria, Jarfalla, SWEDEN
Zhang, Zhiping, Huddinge, SWEDEN

PATENT ASSIGNEE(S): KAROLINSKA INNOVATIONS AB, STOCKHOLM, SWEDEN (non-U.S. corporation)

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 2004241798	A1	20041202
APPLICATION INFO.:	US 2003-619545	A1	20030716 (10)
RELATED APPLN. INFO.:	Continuation of Ser. No. US 2001-720200, filed on 24 May 2001, ABANDONED A 371 of International Ser. No. WO 1999-SE1136, filed on 23 Jun 1999, UNKNOWN		

	NUMBER	DATE
PRIORITY INFORMATION:	SE 1998-2294	19980626
	US 1998-90925P	19980626 (60)
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	APPLICATION	
LEGAL REPRESENTATIVE:	YOUNG & THOMPSON, 745 SOUTH 23RD STREET, 2ND FLOOR, ARLINGTON, VA, 22202	
NUMBER OF CLAIMS:	14	
EXEMPLARY CLAIM:	1	
NUMBER OF DRAWINGS:	6 Drawing Page(s)	
LINE COUNT:	990	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Isolated human nucleic acids implicated in Parkinson's disease and the uses thereof, such as in diagnostic and prognostic methods and in pharmaceutical preparations.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L5 ANSWER 25 OF 117 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN DUPLICATE 1

ACCESSION NUMBER: 2005:104854 BIOSIS

DOCUMENT NUMBER: PREV200500108762

TITLE: A rare truncating mutation in ADH1C (G78Stop) shows significant association with Parkinson disease in a large international sample.

AUTHOR(S): Buervenich, Silvia [Reprint Author]; Carmine, Andrea; Galter, Dagmar; Shahabi, Haydeh N.; Johnels, Bo; Holmberg, Bjorn; Ahlberg, Jarl; Nissbrandt, Hans; Eerola, Johanna; Hellstrom, Olli; Tienari, Pentti J.; Matsuura, Tohru; Ashizawa, Tetsuo; Wullner, Ullrich; Klockgether, Thomas; Zimprich, Alexander; Gasser, Thomas; Hanson, Melissa;

Waseem, Shamaila; Singleton, Andrew; McMahon, Francis J.;
 Anvret, Maria; Sydow, Olof; Olson, Lars
 CORPORATE SOURCE: Mood and Anxiety Disorders Program Genet Unit NIH, Dept Hlth
 and Human Serv, NIMH, 35 Convent Dr, Room 1A-209, Bethesda,
 MD, 20892, USA
 buervens@intra.nimh.nih.gov
 SOURCE: Archives of Neurology, (January 2005) Vol. 62, No. 1, pp.
 74-78. print.
 ISSN: 0003-9942 (ISSN print).
 DOCUMENT TYPE: Article
 LANGUAGE: English
 ENTRY DATE: Entered STN: 16 Mar 2005
 Last Updated on STN: 16 Mar 2005
 AB Background: Alcohol dehydrogenases (ADHs) may be involved in the
 pathogenesis of neurodegenerative disorders because of their multiple
 roles in detoxification pathways and retinoic acid synthesis. In a
 previous study, significant association of an ADH class IV allele
 with Parkinson disease (PD) was found in a Swedish sample.
 Patients: The previously associated single-nucleotide polymorphism
 plus 12 further polymorphisms in the ADH cluster on human
 chromosome 4q23 were screened for association in an extension of the
 original sample that now included 123 Swedish PD patients and 127
 geographically matched control subjects. A rare nonsense
 single-nucleotide polymorphism in ADH1C (G78stop, rs283413) was
 identified in 3 of these patients but in no controls. To obtain
 sufficient power to detect a possible association of this rare variant
 with disease, we screened a large international sample of 1076 PD patients
 of European ancestry and 940 matched controls. Results: The previously
 identified association with an ADH class IV allele remained
 significant ($P < .02$) in the extended Swedish study. Furthermore, in the
 international collaboration, the G78stop mutation in ADH1C was found in 22
 (2.0%) of the PD patients but only in 6 controls (0.6%). This association
 was statistically significant ($\chi^2 = 7.5$; 2-sided $P = .007$; odds ratio, 3.25
 (95% confidence interval, 1.31-8.05)). In addition, the G78stop mutation
 was identified in 4 (10.0%) of 40 Caucasian index cases with PD with
 mainly hereditary forms of the disorder. Conclusion: Findings presented
 herein provide further evidence for mutations in genes encoding ADHs as
 genetic risk factors for PD.

L5 ANSWER 9 OF 117 USPATFULL on STN
 ACCESSION NUMBER: 2006:43224 USPATFULL
 TITLE: Cloning of a gene mutation for parkinson's
 disease.
 INVENTOR(S): Polymeropoulos, Mihael H., Potomac, MD, UNITED STATES
 Lavedan, Christian, North Potomac, MD, UNITED STATES
 Leroy, Elisabeth, Washington, DC, UNITED STATES
 Nussbaum, Robert L., Chevy Chase, MD, UNITED STATES
 Johnson, William G., Short Hills, NJ, UNITED STATES
 Duvoisin, Roger C., Sante Fe, NM, UNITED STATES
 PATENT ASSIGNEE(S): The United States of America as represented by the
 Secretary of the Department of Health and Human
 Services, Washington, DC, UNITED STATES (U.S.
 corporation)

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 7001720	B1	20060221
	WO 9859050		19981230
APPLICATION INFO.:	US 1999-446628		19980625 (9)
	WO 1998-US13071		19980625
			20010919 PCT 371 date

	NUMBER	DATE
PRIORITY INFORMATION:	US 1997-50684P	19970625 (60)

DOCUMENT TYPE: Utility
FILE SEGMENT: GRANTED
PRIMARY EXAMINER: Spector, Lorraine
ASSISTANT EXAMINER: Lockard, Jon M
LEGAL REPRESENTATIVE: Venable LLP, Hobbs, Ann S.
NUMBER OF CLAIMS: 6
EXEMPLARY CLAIM: 1
NUMBER OF DRAWINGS: 11 Drawing Figure(s); 16 Drawing Page(s)
LINE COUNT: 1638

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB **Parkinson's** disease (PD) is a common neurodegenerative disorder with a lifetime incidence of approximately 2 percent. It was recently reported that a PD susceptibility gene is located on the long arm of human chromosome four. The present invention reports the subsequent identification of a mutation in the alpha synuclein gene, which codes for a presynaptic protein thought to be involved in neuronal plasticity. The finding of a specific molecular alteration which is causative for PD will permit the detailed understanding of the pathophysiology of the disorder, which will lead to potential therapeutic interventions, as well as a means for diagnosing individuals having an increased risk of developing the disease.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

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(FILE 'HOME' ENTERED AT 11:20:59 ON 01 MAY 2006)

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 11:21:25 ON 01 MAY 2006
SEA ADH7? OR (ALCOHOL?(S)DEHYDROGENAS?(S) (HUMA? OR SAPIEN?))

7 FILE ADISCTI
5 FILE ADISINSIGHT
3 FILE ADISNEWS
33 FILE AGRICOLA
33 FILE ANABSTR
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9 FILE AQUASCI
34 FILE BIOENG
1281 FILE BIOSIS
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57 FILE DDFU
2317 FILE DGENE
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78 FILE DRUGU
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518 FILE EMBASE
440 FILE ESBIOWASE
10 FILE FROSTI
15 FILE FSTA

1688 FILE GENBANK
 7 FILE HEALSAFE
 44 FILE IFIPAT
 54 FILE JICST-EPLUS
 6 FILE KOSMET
 364 FILE LIFESCI
 600 FILE MEDLINE
 7 FILE NTIS
 2 FILE OCEAN
 424 FILE PASCAL
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 3 FILE RDISCLOSURE
 591 FILE SCISEARCH
 955 FILE TOXCENTER
 886 FILE USPATFULL
 69 FILE USPAT2
 1 FILE WATER
 91 FILE WPIDS
 1 FILE WPIFV
 91 FILE WPINDEX
 8 FILE IPA
 1 FILE NAPRALERT
 3 FILE NLDB

L1 QUE ADH7? OR (ALCOHOL?(S) DEHYDROGENAS?(S) (HUMA? OR SAPIEN?))

 D RANK

FILE 'BIOSIS, TOXCENTER, USPATFULL, CAPLUS, MEDLINE, SCISEARCH, EMBASE, BIOTECHNO, ESBIODBASE, PASCAL, LIFESCI' ENTERED AT 11:25:52 ON 01 MAY 2006

L2 7283 SEA ADH7? OR (ALCOHOL?(S) DEHYDROGENAS?(S) (HUMA? OR SAPIEN?))

L3 1582 SEA L2 AND (ALLELE? OR POLYMORPH?)

L4 127 SEA L3 AND PARKINSON?

L5 117 DUP REM L4 (10 DUPLICATES REMOVED)
 D TI L5 1-117
 D IBIB ABS L5 107 105 99 80 27 25 9

FILE HOME

FILE STNINDEX

FILE BIOSIS

FILE COVERS 1969 TO DATE.

CAS REGISTRY NUMBERS AND CHEMICAL NAMES (CNs) PRESENT
 FROM JANUARY 1969 TO DATE.

RECORDS LAST ADDED: 26 April 2006 (20060426/ED)

FILE TOXCENTER

FILE COVERS 1907 TO 25 Apr 2006 (20060425/ED)

This file contains CAS Registry Numbers for easy and accurate substance identification.

The MEDLINE file segment has been updated with 2006 MEDLINE data and features. See HELP RLOAD for details.

TOXCENTER thesauri in the /CN, /CT, and /MN fields incorporate the MeSH 2006 vocabulary.